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Application Serial Number: 10/612,466
Source: O T P E
Date Processed by STIC: 8/11/2003

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/612,466

DATE: 08/11/2003
TIME: 13:38:59

Input Set : A:\1625seq.001
Output Set : N:\CRF4\08112003\J612466.raw

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66 Asn Ser Asn Ser Thr Glu Phe Val Ser Leu Ala Ser Lys Val Lys Asp
67                               110                               115
69 gcg ctg aag ctg ctg tac agc gga gtc cca ttc ctg gcc ccc tac cac 436
70 Ala Leu Lys Leu Leu Tyr Ser Gly Val Pro Phe Leu Gly Pro Tyr His
71                               125                               130
73 aag gag tgc gct gta acg gcc ttc agc gag gcc agc gtc atc gcc tac 484
74 Lys Glu Ser Ala Val Thr Ala Phe Ser Glu Gly Ser Val Ile Ala Tyr
75                               140                               145
77 tac tgg tct gag ttc agc atc ccg cag cac ctg gta gag gcc gag 532
78 Tyr Trp Ser Glu Phe Ser Ile Pro Gln His Leu Val Glu Glu Ala Glu
79 155                               160                               165
81 cgc gtc atg gcc gag gag cgc gta gtc atg ctg ccc ccg cgg cgc cgc 580
82 Arg Val Met Ala Glu Glu Arg Val Val Met Leu Pro Pro Arg Ala Arg
83                               175                               180
85 tcc ctg aag tcc ttt gta gtc acc tca gta gta gct ttc ccc acc gac 628
86 Ser Leu Lys Ser Phe Val Val Thr Ser Val Val Ala Phe Pro Thr Asp
87                               190                               195
89 tcc aaa aca gta cag agc acc cag gac aac agc tgc agc ttt gcc ctg 676
90 Ser Lys Thr Val Gln Arg Thr Gln Asp Asn Ser Cys Ser Phe Gly Leu
91                               205                               210
93 cac gcc cgc ggt gta gag ctg atg cgc ttc acc acg ccc gcc ttc cct 724
94 His Ala Arg Gly Val Glu Leu Met Arg Phe Thr Thr Pro Gly Phe Pro
95                               220                               225
97 gag agc ccc tac ccc gct cat gcc cgc tgc cag tgg gcc ctg cgg ggg 772
98 Asp Ser Pro Tyr Pro Ala His Ala Arg Cys Gln Trp Ala Leu Arg Gly
99 235                               240                               245
101 gag gcc gac tca gta gta gta gta gta gta gta gta gta gta gta 820
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103                               255                               260
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106 Ser Cys Asp Glu Arg Gly Ser Asp Leu Val Thr Val Tyr Asn Thr Leu
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110 Ser Pro Met Glu Pro His Ala Leu Val Gln Leu Cys Gly Thr Tyr Pro
111                               285                               290
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114 Pro Ser Tyr Asn Leu Thr Phe His Ser Ser Gln Asn Val Leu Leu Ile
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121 ttc ttc cag ctg cct agc atg agc agc tgt gga gcc cgc tta cgt aaa 1060
122 Phe Phe Gln Leu Pro Arg Met Ser Ser Cys Gly Gly Arg Leu Arg Lys
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Input Set : A:\1625seq.001

Output Set : N:\CRF4\08112003\J612466.raw

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133	aag gtg agc ttc aaa ttc ttc tac ctg ctg gag ccc ggc gtg cct gcg			1204
134	Lys Val Ser Phe Lys Phe Phe Tyr Leu Leu Glu Pro Gly Val Pro Ala			
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137	ggc acc tgc ccc aag gac tac gtg gag atc aat ggg gag aaa tac tgc			1252
138	Gly Thr Cys Pro Lys Asp Tyr Val Glu Ile Asn Gly Glu Lys Tyr Cys			
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141	gga gag agg tcc cag ttc gtc gtc acc agc aac agc aac aag atc aca			1300
142	Gly Glu Arg Ser Gln Phe Val Val Thr Ser Asn Ser Asn Lys Ile Thr			
143	415	420	425	
145	gtt cgc ttc cac tca gat cag tcc tac acc gac acc ggc ttc tta gct			1348
146	Val Arg Phe His Ser Asp Gln Ser Tyr Thr Asp Thr Gly Phe Leu Ala			
147	430	435	440	
149	gaa tac ctc tcc tac gac tcc agt gac cca tgc ccg ggc cag ttc acg			1396
150	Glu Tyr Leu Ser Tyr Asp Ser Ser Asp Pro Cys Pro Gly Gln Phe Thr			
151	445	450	455	
153	tgc cgc acg ggc cgg tgt atc cgg aag gag ctg cgc tgt gat ggc tgg			1444
154	Cys Arg Thr Gly Arg Cys Ile Arg Lys Glu Leu Arg Cys Asp Gly Trp			
155	460	465	470	
157	gcc cag tgc acc gac cac agc gat gag ctc aac tgc agt tgc gac gcc			1492
158	Ala Asp Cys Thr Asp His Ser Asp Glu Leu Asn Cys Ser Cys Asp Ala			
159	475	480	485	
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162	Gly His Gln Phe Thr Cys Lys Asn Lys Phe Cys Lys Pro Leu Phe Trp			
163	495	500	505	
165	gtc tgc gac agt gtg aac gac tgc gga gac aac agc gac gag cag ggc			1588
166	Val Cys Asp Ser Val Asn Asp Cys Gly Asp Asn Ser Asp Glu Gln Gly			
167	510	515	520	
169	tgc agt tgt ccg gcc cag acc ttc agg tgt tcc aat ggg aag tgc ctc			1636
170	Cys Ser Cys Pro Ala Gln Thr Phe Arg Cys Ser Asn Gly Lys Cys Leu			
171	525	530	535	
173	tcg aaa agc cag cag tgc aat ggg aag gac gac tgt ggg gac ggc tcc			1684
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182	Thr Tyr Arg Cys Leu Asn Gly Leu Cys Lys Ser Lys Gly Asn Pro Glu			
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185	tgt gac ggc aag gag gac tgt agc gac ggc tca gat gag aag gac tgc			1828
186	Cys Asp Gly Lys Glu Asp Cys Ser Asp Gly Ser Asp Glu Lys Asp Cys			
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189	gac tgt ggc ctg cgg tca ttc acg aga cag gct cgt gtt gtt ggg ggc			1876
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Input Set : A:\1625seq.001

Output Set: N:\CRF4\08112003\J612466.raw

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197 ctg ggc cag ggc cac atc tgc ggt gct tcc ctc atc tct ccc aac tgg 1972
198 Leu Gly Gln Gly His Ile Cys Gly Ala Ser Leu Ile Ser Pro Asn Trp
199 635      640      645
201 ctg gtc tct gcc gca cac tgc tac atc gat gac aga gga ttc agg tac 2020
202 Leu Val Ser Ala Ala His Cys Tyr Ile Asp Asp Arg Gly Phe Arg Tyr
203      655      660      665
205 tca gac ccc acg cag tgg acg gcc ttc ctg ggc ttc cag cag agc 2068
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209 cag cgc agc gcc cct ggg gtg cag gag cgc agg ctc aag cgc atc atc 2116
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211      685      690      695
213 tcc cac ccc ttc ttc aat gac ttc acc ttc gac tat gac atc gcg ctg 2164
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217 ctg gag ctg gag aaa ccg gca gag tac agc tcc atg gtg cgg ccc atc 2212
218 Leu Glu Leu Glu Lys Pro Ala Glu Tyr Ser Ser Met Val Arg Pro Ile
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222 Cys Leu Pro Asp Ala Ser His Val Phe Pro Ala Gly Lys Ala Ile Trp
223      735      740      745
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226 Val Thr Gly Trp Gly His Thr Gln Tyr Gly Gly Thr Gly Ala Leu Ile
227      750      755      760
229 ctg caa aag ggt gag atc cgc gtc atc aac cag acc acc tgc gag aac 2356
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233 etc ctg ccg cag cag atc acg ccg cgc atg atg tgc gtg ggc ttc ctc 2404
234 Leu Leu Pro Gln Gln Ile Thr Pro Arg Met Met Cys Val Gly Phe Leu
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237 agc ggc ggc gtg gac tcc tgc cag ggt gat tcc ggg gga ccc ctg tcc 2452
238 Ser Gly Gly Val Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Ser
239 795      800      805      810
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242 Ser Val Glu Ala Asp Gly Arg Ile Phe Gln Ala Gly Val Val Ser Trp
243      815      820      825      830
245 gga gac ggc tgc gct cag agg aac aag cca ggc gtg tac aca agg ctc 2548
246 Gly Asp Gly Cys Ala Gln Arg Asn Lys Pro Gly Val Tyr Thr Arg Leu
247      835      840
249 cct ctg ttt cgg gac tgg atc aas gag aac act ggg gta ta ggggcccggg 2599
250 Pro Leu Phe Arg Asp Trp Ile Lys Glu Asn Thr Gly Val
251      845      850      855
253 ccacccaat gtgtacacct gcggggccac ccacgtgtca cccagtggtg cagccttca 2659
254 gctggagac tggaccgctg actgcaccag cgcgccaga acatacactg tgaactcaat 2719
255 ctccagggtc ccaaatctgc ctageaaacc tctcgcttc tcagcctcca aagtggagct 2779
256 gggaggtaga agggaggagc actggtggtt ctactgacc aactgggggc aaaggtttga 2839

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/612,466

DATE: 08/11/2003
TIME: 13:38:59

Input Set : A:\1625seq.001
Output Set: N:\CRF4\08112003\J612466.raw

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259 ggctgcggga tctgggtgt ggggcccttg ggcacagctc ttgaggagc ccaggctcgg 3019
260 aggacccctg aaaaacagac ggtctgagac tgaattgtt ttaccagctc ccagggtgga 3079
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267 <213> ORGANISM: Homo Sapien
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274 Glu Glu Gly Val Glu Phe Leu Pro Val Asn Asn Val Lys Lys Val Glu
275 35 40 45
276 Lys His Gly Pro Gly Arg Trp Val Val Leu Ala Val Leu Ile Gly
277 50 55 60
278 Leu Leu Leu Val Leu Leu Gly Ile Gly Phe Leu Val Trp His Leu Gln
279 65 70 75 80
280 Tyr Arg Asp Val Arg Val Gln Lys Val Phe Asn Gly Tyr Met Arg Ile
281 85 90 95
282 Thr Asn Glu Asn Phe Val Asp Ala Tyr Glu Asn Ser Asn Ser Thr Glu
283 100 105 110
284 Phe Val Ser Leu Ala Ser Lys Val Lys Asp Ala Leu Lys Leu Leu Tyr
285 115 120 125
286 Ser Gly Val Pro Phe Leu Gly Pro Tyr His Lys Glu Ser Ala Val Thr
287 130 135 140
288 Ala Phe Ser Glu Gly Ser Val Ile Ala Tyr Tyr Trp Ser Glu Phe Ser
289 145 150 155 160
290 Ile Pro Gln His Leu Val Glu Glu Ala Glu Arg Val Met Ala Glu Glu
291 165 170 175
292 Arg Val Val Met Leu Pro Pro Arg Ala Arg Ser Leu Lys Ser Phe Val
293 180 185 190
294 Val Thr Ser Val Val Ala Phe Pro Thr Asp Ser Lys Thr Val Gln Arg
295 195 200 205
296 Thr Gln Asp Asn Ser Cys Ser Phe Gly Leu His Ala Arg Gly Val Glu
297 210 215 220
298 Leu Met Arg Phe Thr Thr Pro Gly Phe Pro Asp Ser Pro Tyr Pro Ala
299 225 230 235 240
300 His Ala Arg Cys Gln Trp Ala Leu Arg Gly Asp Ala Asp Ser Val Leu
301 245 250 255
302 Ser Leu Thr Phe Arg Ser Phe Asp Leu Ala Ser Cys Asp Gly Arg Gly
303 260 265 270
304 Ser Asp Leu Val Thr Val Tyr Asn Thr Leu Ser Pro Met Glu Pro His
305 275 280 285
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10/6/246

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Pro Ser Ala Leu Pro Ile Cys Leu His Pro Ala Gly Ile Pro Pro Gly
35 40 45
Ala Ser Cys Trp Val Leu Gly Trp Lys Glu Pro Gln Asp Arg Val Pro
50 55 60
Val Ala Ala Ala Val Ser Ile Leu Thr Gln Arg Ile Cys Asp Cys Leu
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20 25 30
Phe Phe Pro Leu Gln
35

which response is the valid one?

If it is 'Artificial Sequence' include (220)-(223)

*section
and explanation
of source
material on
(223) line*

same error